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# SEQUENCE LISTING

<110> RHODES, Simon J.  
BRIDWELL, Jeanne L.  
MEIER, Bradley C.  
PARKER, Gretchen E.  
PRICE, Jeffrey R.  
SHOWALTER, Aaron D.  
SLOOP, Kyle W.

<120> GENERATION OF DIAGNOSTIC TOOLS TO ASSAY THE HUMAN  
LHX3/P-LIM/LIM-3 FACTOR

<130> 053884-5003

<140> NOT YET ASSIGNED

<141> 2001-08-17

<150> PCT/US00/04424

<151> 2000-02-22

<150> US 60/121,110

<151> 1999-02-22

<160> 113

<170> PatentIn Ver. 2.1

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Pro Gln Ser Met Arg Val Leu Ala Gly Asn Gly Pro Ser Ser Asp Leu		
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 Phe Arg Asn Met Lys Arg Ser Arg Gly Asn Ser Lys Ser Asp Lys Asp  
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 245 250 255  
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 260 265 270  
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 Thr Gln Val Val Arg Arg Ala Gln Asp Phe Val Tyr His Leu His Cys  
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 Pro Lys Pro Ala Arg His Val Arg Glu Gln Leu Ser Thr Glu Thr Gly  
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 Glu Lys Arg Leu Lys Lys Asp Ala Gly Arg Gln Arg Trp Gly Gln Tyr  
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 Asn Thr Ser Pro Lys Pro Ala Arg His Val Arg Glu Gln Leu Ser Ser  
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Gln Tyr Arg Glu Leu Arg Pro Gly Ser Pro Tyr Gly Val Pro Pro Ser		
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<210> 12

<211> 402

<212> PRT

<213> Homo sapiens

<400> 12

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Asp Leu Leu Leu Ala Leu Leu Ala Arg Arg Ala Asp Leu Arg Arg Glu
                20                      25                     30

Ile Pro Leu Cys Ala Gly Cys Asp Gln His Ile Leu Asp Arg Phe Ile
    35                      40                     45

Leu Lys Ala Leu Asp Arg His Trp His Ser Lys Cys Leu Lys Cys Ser
    50                      55                     60

Asp Cys His Thr Pro Leu Ala Glu Arg Cys Phe Ser Arg Gly Glu Ser
    65                      70                     75                     80

Val Tyr Cys Lys Asp Asp Phe Phe Lys Arg Phe Gly Thr Lys Cys Ala
                85                      90                     95

Ala Cys Gln Leu Gly Ile Pro Pro Thr Gln Val Val Arg Arg Ala Gln
    100                      105                     110

Asp Phe Val Tyr His Leu His Cys Phe Ala Cys Val Val Cys Lys Arg
    115                      120                     125

Gln Leu Ala Thr Gly Asp Glu Phe Tyr Leu Met Glu Asp Ser Arg Leu
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Val	Cys	Lys	Ala	Asp	Tyr	Glu	Thr	Ala	Lys	Gln	Arg	Glu	Ala	Glu	Ala	
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Glu	Gln	Leu	Ser	Ser	Glu	Thr	Gly	Leu	Asp	Met	Arg	Val	Val	Gln	Val	
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Pro	Ala	Asn	Gly	Leu	Tyr	Gly	Ser	Leu	Gly	Glu	Pro	Thr	Gln	Ala	Leu	
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Leu	Ala	Gly	Pro	Glu	Gln	Tyr	Arg	Glu	Leu	Arg	Pro	Gly	Ser	Pro	Tyr	
305					310					315					320	
Gly	Val	Pro	Pro	Ser	Pro	Ala	Ala	Pro	Gln	Ser	Leu	Pro	Gly	Pro	Gln	
				325					330					335		
Pro	Leu	Leu	Ser	Ser	Leu	Val	Tyr	Pro	Asp	Thr	Ser	Leu	Gly	Leu	Val	
			340					345					350			
Pro	Ser	Gly	Ala	Pro	Gly	Gly	Pro	Pro	Pro	Met	Arg	Val	Leu	Ala	Gly	
		355					360					365				
Asn	Gly	Pro	Ser	Ser	Asp	Leu	Ser	Thr	Gly	Ser	Ser	Gly	Gly	Tyr	Pro	
	370					375						380				
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Gln Phe

<210> 13  
 <211> 1658  
 <212> DNA  
 <213> Sus scrofa

<400> 13

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tgcagtgact gccacacgcc gctggccgag cgctgcttca gccgcggaga gagcctctac 240  
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aagcggccgc gcacgaccat cacggccaag cagctggaga cgctgaagag cgcctacaac 540  
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gacgccggcc ggcagcgctg gggccagtac tttcgttaaca tgaagcgcgc ccgcggtggc 720  
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<210> 14

<211> 401

<212> PRT

<213> Sus scrofa

<400> 14

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Leu Cys Ala Gly Cys Asp Gln His Ile Leu Asp Arg Phe Ile Leu Lys  
35 40 45  
Ala Leu Asp Arg His Trp His Ser Lys Cys Leu Lys Cys Ser Asp Cys  
50 55 60  
His Thr Pro Leu Ala Glu Arg Cys Phe Ser Arg Gly Glu Ser Leu Tyr  
65 70 75 80  
Cys Lys Asp Asp Phe Phe Lys Arg Phe Gly Thr Lys Cys Ala Ala Cys  
85 90 95  
Gln Leu Gly Ile Pro Pro Thr Gln Val Val Arg Arg Ala Gln Asp Phe  
100 105 110

Val Tyr His Leu His Cys Phe Ala Cys Val Val Cys Lys Arg Gln Leu  
 115 120 125  
 Ala Thr Gly Asp Glu Phe Tyr Leu Met Glu Asp Ser Arg Leu Val Cys  
 130 135 140  
 Lys Ala Asp Tyr Glu Thr Ala Lys Gln Arg Glu Ala Glu Ala Thr Ala  
 145 150 155 160  
 Lys Arg Pro Arg Thr Thr Ile Thr Ala Lys Gln Leu Glu Thr Leu Lys  
 165 170 175  
 Ser Ala Tyr Asn Thr Ser Pro Lys Pro Ala Arg His Val Arg Glu Gln  
 180 185 190  
 Leu Ser Ser Glu Thr Gly Leu Asp Met Arg Val Val Gln Val Trp Phe  
 195 200 205  
 Gln Asn Arg Arg Ala Lys Glu Lys Arg Leu Lys Lys Asp Ala Gly Arg  
 210 215 220  
 Gln Arg Trp Gly Gln Tyr Phe Arg Asn Met Lys Arg Ala Arg Gly Gly  
 225 230 235 240  
 Ser Lys Ser Asp Lys Asp Ser Val Gln Glu Glu Gly Gln Asp Ser Asp  
 245 250 255  
 Ala Glu Val Ser Phe Thr Asp Glu Pro Ser Met Ala Glu Met Gly Pro  
 260 265 270  
 Ala Asn Gly Leu Tyr Gly Gly Leu Gly Glu Pro Ala Pro Ala Leu Gly  
 275 280 285  
 Arg Pro Ser Gly Ala Pro Gly Ser Phe Pro Leu Glu His Gly Gly Leu  
 290 295 300  
 Ala Gly Pro Glu Gln Tyr Gly Glu Leu Arg Pro Ser Ser Pro Tyr Gly  
 305 310 315 320  
 Val Pro Ser Ser Pro Ala Ala Leu Gln Ser Leu Pro Gly Pro Gln Pro  
 325 330 335  
 Leu Leu Ser Ser Leu Val Tyr Pro Glu Ala Gly Leu Gly Leu Val Pro  
 340 345 350  
 Ala Gly Pro Pro Gly Gly Pro Pro Pro Met Arg Val Leu Ala Gly Asn  
 355 360 365  
 Gly Pro Ser Ser Asp Leu Ser Thr Gly Ser Ser Gly Gly Tyr Pro Asp  
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 Phe Pro Ala Ser Pro Ala Ser Trp Leu Asp Glu Val Asp His Ala Gln  
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 Phe

<210> 15  
 <211> 1664  
 <212> DNA  
 <213> *Sus scrofa*

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 aagaaggacg ccggccggca gcgtcggggc cagtactttc gtaacatgaa gcgcgccccg 720  
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 ctgctctttc tagaccggag tggtcagccc ccgaagccgg ggagggggggc tctccccagc 1560  
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<210> 16  
 <211> 403  
 <212> PRT  
 <213> *Sus scrofa*

<400> 16  
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 Ile Pro Leu Cys Ala Gly Cys Asp Gln His Ile Leu Asp Arg Phe Ile  
 35 40 45  
 Leu Lys Ala Leu Asp Arg His Trp His Ser Lys Cys Leu Lys Cys Ser  
 50 55 60  
 Asp Cys His Thr Pro Leu Ala Glu Arg Cys Phe Ser Arg Gly Glu Ser

65	70	75	80
Leu Tyr Cys Lys Asp	Asp Phe Phe Lys Arg	Phe Gly Thr Lys Cys Ala	
	85	90	95
Ala Cys Gln Leu Gly Ile Pro Pro Thr Gln Val Val Arg Arg Ala Gln			
	100	105	110
Asp Phe Val Tyr His Leu His Cys Phe Ala Cys Val Val Cys Lys Arg			
	115	120	125
Gln Leu Ala Thr Gly Asp Glu Phe Tyr Leu Met Glu Asp Ser Arg Leu			
	130	135	140
Val Cys Lys Ala Asp Tyr Glu Thr Ala Lys Gln Arg Glu Ala Glu Ala			
	145	150	155
Thr Ala Lys Arg Pro Arg Thr Thr Ile Thr Ala Lys Gln Leu Glu Thr			
	165	170	175
Leu Lys Ser Ala Tyr Asn Thr Ser Pro Lys Pro Ala Arg His Val Arg			
	180	185	190
Glu Gln Leu Ser Ser Glu Thr Gly Leu Asp Met Arg Val Val Gln Val			
	195	200	205
Trp Phe Gln Asn Arg Arg Ala Lys Glu Lys Arg Leu Lys Lys Asp Ala			
	210	215	220
Gly Arg Gln Arg Trp Gly Gln Tyr Phe Arg Asn Met Lys Arg Ala Arg			
	225	230	235
Gly Gly Ser Lys Ser Asp Lys Asp Ser Val Gln Glu Glu Gly Gln Asp			
	245	250	255
Ser Asp Ala Glu Val Ser Phe Thr Asp Glu Pro Ser Met Ala Glu Met			
	260	265	270
Gly Pro Ala Asn Gly Leu Tyr Gly Gly Leu Gly Glu Pro Ala Pro Ala			
	275	280	285
Leu Gly Arg Pro Ser Gly Ala Pro Gly Ser Phe Pro Leu Glu His Gly			
	290	295	300
Gly Leu Ala Gly Pro Glu Gln Tyr Gly Glu Leu Arg Pro Ser Ser Pro			
	305	310	315
Tyr Gly Val Pro Ser Ser Pro Ala Ala Leu Gln Ser Leu Pro Gly Pro			
	325	330	335
Gln Pro Leu Leu Ser Ser Leu Val Tyr Pro Glu Ala Gly Leu Gly Leu			
	340	345	350
Val Pro Ala Gly Pro Pro Gly Gly Pro Pro Pro Met Arg Val Leu Ala			
	355	360	365
Gly Asn Gly Pro Ser Ser Asp Leu Ser Thr Gly Ser Ser Gly Gly Tyr			

370 375 380  
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 385 390 395 400

Ala Gln Phe

<210> 17  
 <400> 17  
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<210> 18  
 <400> 18  
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<210> 19  
 <211> 440  
 <212> PRT  
 <213> *Drosophila melanogaster*

<400> 19  
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 Glu Phe Leu Leu Ser Thr Ile Pro Lys Cys Gly Gly Cys His Glu Leu  
 35 40 45  
 Ile Leu Asp Arg Phe Ile Leu Lys Val Leu Glu Arg Thr Trp His Ala  
 50 55 60  
 Lys Cys Leu Gln Cys Ser Glu Cys His Gly Gln Leu Asn Asp Lys Cys  
 65 70 75 80  
 Phe Ala Arg Asn Gly Gln Leu Phe Cys Lys Glu Asp Phe Phe Lys Arg  
 85 90 95  
 Tyr Gly Thr Lys Cys Ser Ala Cys Asp Met Gly Ile Pro Pro Thr Gln  
 100 105 110  
 Val Val Arg Arg Ala Gln Asp Asn Val Tyr His Leu Gln Cys Phe Leu  
 115 120 125  
 Cys Ala Met Cys Ser Arg Thr Leu Asn Thr Gly Asp Glu Phe Tyr Leu  
 130 135 140  
 Met Glu Asp Arg Lys Leu Ile Cys Lys Arg Asp Tyr Glu Glu Ala Lys  
 145 150 155 160  
 Ala Lys Gly Leu Tyr Leu Asp Gly Ser Leu Asp Gly Asp Gln Pro Asn  
 165 170 175



<400> 20

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Lys Phe Ile Leu Lys Val Leu Asp Arg His Trp His Ser Ser Cys Leu  
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Lys Cys Ala Asp Cys Gln Met Gln Leu Ala Asp Arg Cys Phe Ser Arg  
35 40 45

Ala Gly Ser Val Tyr Cys Lys Glu Asp Phe Phe Lys Arg Phe Gly Thr  
50 55 60

Lys Cys Thr Ala Cys Gln Gln Gly Ile Pro Pro Thr Gln Val Val Arg  
65 70 75 80

Lys Ala Gln Asp Phe Val Tyr His Leu His Cys Phe Ala Cys Ile Ile  
85 90 95

Cys Asn Arg Gln Leu Ala Thr Gly Asp Glu Phe Tyr Leu Met Glu Asp  
100 105 110

Gly Arg Leu Val Cys Lys Glu Asp Tyr Glu Thr Ala Lys Gln Asn Asp  
115 120 125

Asp Ser Glu Ala Gly Ala Lys Arg Pro Arg Thr Thr Ile Thr Ala Lys  
130 135 140

Gln Leu Glu Thr Leu Lys Asn Ala Tyr Lys Asn Ser Pro Lys Pro Ala  
145 150 155 160

Arg His Val Arg Glu Gln Leu Ser Ser Glu Thr Gly Leu Asp Met Arg  
165 170 175

Val Val Gln Val Trp Phe Gln Asn Arg Arg Ala Lys Glu Lys Arg Leu  
180 185 190

Lys Lys Asp Ala Gly Arg His Arg Trp Gly Gln Phe Tyr Lys Ser Val  
195 200 205

Lys Arg Ser Arg Gly Gly Ser Lys Gln Glu Lys Glu Ser Ser Ala Glu  
210 215 220

Asp Cys Gly Val Ser Asp Ser Glu Leu Ser Phe Arg Glu Asp Gln Ile  
225 230 235 240

Leu Ser Glu Leu Gly His Thr Asn Arg Ile Tyr Gly Asn Val Gly Asp  
245 250 255

Val Thr Gly Gly Gln Leu Met Asn Gly Ser Phe Ser Met Asp Gly Thr  
260 265 270

Gly Gln Ser Tyr Gln Asp Leu Arg Asp Gly Ser Pro Tyr Gly Ile Pro  
275 280 285

Gln Ser Pro Ser Ser Ile Ser Ser Leu Pro Ser His Ala Pro Leu Leu

290	295	300
Asn Gly Leu Asp Tyr Thr Val Asp Ser Asn Leu Gly Ile Ile Ala His		
305	310	315 320
Ala Gly Gln Gly Val Ser Gln Thr Leu Arg Ala Met Ala Gly Gly Pro		
	325	330 335
Thr Ser Asp Leu Ser Thr Gly Ser Ser Val Gly Tyr Pro Asp Phe Pro		
	340	345 350
Thr Ser Pro Ala Ser Trp Leu Asp Glu Met Asp His Pro Pro Phe		
	355	360 365

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 <211> 402  
 <212> PRT  
 <213> Mus musculus

<400> 21  
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 Ile Pro Met Cys Ala Gly Cys Asp Gln His Ile Leu Asp Arg Phe Ile  
 35 40 45  
 Leu Lys Ala Leu Asp Arg His Trp His Ser Lys Cys Leu Lys Cys Ser  
 50 55 60  
 Asp Cys His Val Pro Leu Ala Glu Arg Cys Phe Ser Arg Gly Glu Ser  
 65 70 75 80  
 Val Tyr Cys Lys Asp Asp Phe Phe Lys Arg Phe Gly Thr Lys Cys Ala  
 85 90 95  
 Ala Cys Gln Leu Gly Ile Pro Pro Thr Gln Val Val Arg Arg Ala Gln  
 100 105 110  
 Asp Phe Val Tyr His Leu His Cys Phe Ala Cys Val Val Cys Lys Arg  
 115 120 125  
 Gln Leu Ala Thr Gly Asp Glu Phe Tyr Leu Met Glu Asp Ser Arg Leu  
 130 135 140  
 Val Cys Lys Ala Asp Tyr Glu Thr Ala Lys Gln Arg Glu Ala Glu Ala  
 145 150 155 160  
 Thr Ala Lys Arg Pro Arg Thr Thr Ile Thr Ala Lys Gln Leu Glu Thr  
 165 170 175  
 Leu Lys Ser Ala Tyr Asn Thr Ser Pro Lys Pro Ala Arg His Val Arg  
 180 185 190

Glu Gln Leu Ser Ser Glu Thr Gly Leu Asp Met Arg Val Val Gln Val  
 195 200 205  
 Trp Phe Gln Asn Arg Arg Ala Lys Glu Lys Arg Leu Lys Lys Asp Ala  
 210 215 220  
 Gly Arg Gln Arg Trp Gly Gln Tyr Phe Arg Asn Met Lys Arg Ser Arg  
 225 230 235 240  
 Gly Ser Ser Lys Ser Asp Lys Asp Ser Ile Gln Glu Gly Gln Asp Ser  
 245 250 255  
 Asp Ala Glu Val Ser Phe Thr Asp Glu Pro Ser Met Ala Asp Met Gly  
 260 265 270  
 Pro Ala Asn Gly Leu Tyr Ser Ser Leu Gly Glu Pro Ala Pro Ala Leu  
 275 280 285  
 Gly Arg Pro Val Gly Gly Leu Gly Ser Phe Thr Leu Asp His Gly Gly  
 290 295 300  
 Leu Thr Gly Pro Glu Gln Tyr Arg Glu Leu Arg Pro Gly Ser Pro Tyr  
 305 310 315 320  
 Gly Ile Pro Pro Ser Pro Ala Ala Pro Gln Ser Leu Pro Gly Pro Gln  
 325 330 335  
 Pro Leu Leu Ser Ser Leu Val Tyr Pro Asp Thr Asn Leu Ser Leu Val  
 340 345 350  
 Pro Ser Gly Pro Pro Gly Gly Pro Pro Pro Met Arg Val Leu Ala Gly  
 355 360 365  
 Asn Gly Pro Ser Ser Asp Leu Ser Thr Glu Ser Ser Ser Gly Tyr Pro  
 370 375 380  
 Asp Phe Pro Ala Ser Pro Ala Ser Trp Leu Asp Glu Val Asp His Ala  
 385 390 395 400  
 Gln Phe

<210> 22

<211> 8867

<212> DNA

<213> Homo sapiens

<400> 22

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<211> 2070

<212> DNA

<213> Homo sapiens

<400> 27

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<210> 28
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Ala Ala Val Cys Thr Leu Gly Gly Thr Arg
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<210> 29
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Asp Leu Leu Leu Ala Leu Leu Ala Arg Arg Ala Asp Leu Arg Arg
          20             25             30

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<210> 31

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1 5 10 15  
Pro Ala Ala Ala Ala Val Cys Thr Leu Pro Gly Thr Arg  
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<210> 32  
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<210> 33  
<211> 31  
<212> PRT  
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<400> 33  
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1 5 10 15  
Asp Leu Leu Leu Ala Leu Leu Ala Arg Arg Glu Asp Leu Arg Arg  
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<210> 34  
<211> 22  
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<220>  
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<400> 34  
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<210> 35  
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<400> 35  
ccgagtcctg cccaaggtgc 20

<210> 36  
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 <210> 40  
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 <400> 40  
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<210> 41  
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 <210> 42  
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 <210> 43  
 <211> 26  
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 binding sequence  
  
 <400> 43  
 gatcccagaa aattaattaa ttgtaa 26  
  
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<400> 45  
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<210> 46  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 46  
tgacctcgga ggagcgcgtc t 21

<210> 47  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 47  
tcgtccttgc agtaaacgct 20

<210> 48  
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<212> DNA  
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<220>  
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<400> 48  
agcgtttact gcaaggacga 20

<210> 49  
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<220>  
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<400> 49  
cgcacttggt cccgaagcgc 20

<210> 50  
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<220>  
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<400> 50  
gcgcttcggg accaagtgcg 20

<210> 51  
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<400> 51  
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<210> 52  
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<400> 52  
ggacaaggac agcgttcag 19

<210> 53  
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<400> 53  
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<210> 54  
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<220>  
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<400> 54  
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<210> 55

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 <210> 58  
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<210> 60

<211> 32

<212> DNA

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<210> 61

<211> 29

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<223> Description of Artificial Sequence:PCR primer

<400> 61

cggaattcag tcagaactga gcgtgatcc 29

<210> 62

<211> 33

<212> DNA

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<223> Description of Artificial Sequence:PCR primer

<400> 62

cgggatccaa gcagcgagag gccgaggcca cgg 33

<210> 63

<211> 29

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<223> Description of Artificial Sequence:PCR primer

<400> 63

cggaattcag tcagaactga gcgtgatcc 29

<210> 64

<211> 28

<212> DNA

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<223> Description of Artificial Sequence:PCR primer

<400> 64

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28

<210> 65

<211> 28

<212> DNA

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<223> Description of Artificial Sequence:PCR primer

<400> 65

cacatttaata tagctaagta cctaatagt

28

<210> 66

<211> 28

<212> DNA

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<223> Description of Artificial Sequence:PCR primer

<400> 66

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28

<210> 67

<211> 28

<212> DNA

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<223> Description of Artificial Sequence:PCR primer

<400> 67

cacatttggc gcgccaagta cctaatagt

28

<210> 68

<211> 34

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:PCR primer

<400> 68

cgggatccat gctggatcgg gatgtgggcc caac

34

<210> 69

<211> 32

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:PCR primer

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<210> 70  
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<220>

<223> Description of Artificial Sequence:PCR primer

<400> 70  
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<210> 71  
<211> 27  
<212> DNA  
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<223> Description of Artificial Sequence:PCR primer

<400> 71  
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<210> 72  
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<223> Description of Artificial Sequence:PCR primer

<400> 72  
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<210> 73  
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<223> Description of Artificial Sequence:PCR primer

<400> 73  
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<210> 74  
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<210> 75  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 75  
ggacaaggac agcggttcag 19

<210> 76  
<211> 18  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 76  
ctcccgtaga ggccattg 18

<210> 77  
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<220>  
<223> Description of Artificial Sequence:Probe

<400> 77  
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<210> 78  
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<223> Description of Artificial Sequence:PCR primer

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cggaattcta caacacctcg cccaagccgg

30

<210> 79

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:PCR primer

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27

<210> 80

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:PCR primer

<400> 80

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28

<210> 81

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:PCR primer

<400> 81

cgggatcctt gatatttacc ccggaggc

28

<210> 82

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:PCR primer

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32

<210> 83

<211> 29

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 <210> 84  
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 <223> Description of Artificial Sequence:PCR primer  
  
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 <210> 85  
 <211> 29  
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 <223> Description of Artificial Sequence:PCR primer  
  
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 <210> 86  
 <211> 19  
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 <223> Description of Artificial Sequence:PCR primer  
  
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 <210> 87  
 <211> 18  
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 <223> Description of Artificial Sequence:PCR primer  
  
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 <210> 88  
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<212> DNA  
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<210> 89  
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<210> 90  
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<400> 90  
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<210> 91  
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<220>  
 <223> Description of Artificial Sequence:PCR primer

<400> 91  
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<210> 92  
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<220>  
 <223> Description of Artificial Sequence:PCR primer

<400> 92  
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<210> 93  
<211> 19  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 93  
gaagttcagg gtcggaggg

19

<210> 94  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 94  
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20

<210> 95  
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<400> 95  
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21

<210> 96  
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<220>  
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<400> 96  
cgggatccat gctgctggaa acggggct

28

<210> 97  
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<400> 97  
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<210> 98  
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<220>  
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<210> 99  
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<223> Description of Artificial Sequence:PCR primer

<400> 99  
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<210> 100  
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acattagcta cttagcta taaatgtg 28

<210> 101  
<211> 28  
<212> DNA  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 101  
cacatttaat tagctaagta cctaattg 28

<210> 102  
<211> 192  
<212> DNA  
<213> Homo sapiens

<400> 102  
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 caccttcccc gggagaagct ttccccaatc cccaggtctc tagatcattc tggtctcgag 120  
 tatcctgtgg aggaggcaaa aatgcctggc gcccttctc tccaagctca attctctaag 180  
 cccctcaggg tc 192

<210> 103  
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 <212> DNA  
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<220>  
 <223> Description of Artificial Sequence:PCR primer

<400> 103  
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<210> 104  
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<220>  
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<400> 104  
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<210> 105  
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<220>  
 <223> Description of Artificial Sequence:PCR primer

<400> 105  
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<210> 106  
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 <212> DNA  
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<220>  
 <223> Description of Artificial Sequence:PCR primer

<400> 106  
 tacgaggtga cccagaactt 20

<210> 107  
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<212> DNA  
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<210> 108  
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<210> 109  
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<210> 110  
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<210> 112  
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<210> 113  
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<220>  
<223> Description of Artificial Sequence:PCR primer

<400> 113  
cctcgtgtga ggtgcagggt 20

Sequence